

p7.5tk

7.5 k promoter

Not I

Apa I

GGCCAAAATTGAAAAAAGCTAGATCTATTATTGCACCGGGCCGCGCAATGGGCCCGGCCGCCAACGGCGGA

Met Gly Pro Ala Ala Asn Gly Gly

tk coding sequence

pE/Ltk

synthetic E/L promoter

Not I

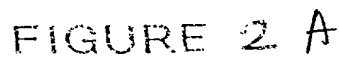
Apa I

GGCCAAAATTGAAATTTTATTTTTTTTGGAAATATAAAGCGGGCCGCGCAATGGGCCCGGCCGCCAACGGCGGA

Met Gly Pro Ala Ala Asn Gly Gly

tk coding sequence

FIGURE 1

[illegible]

B

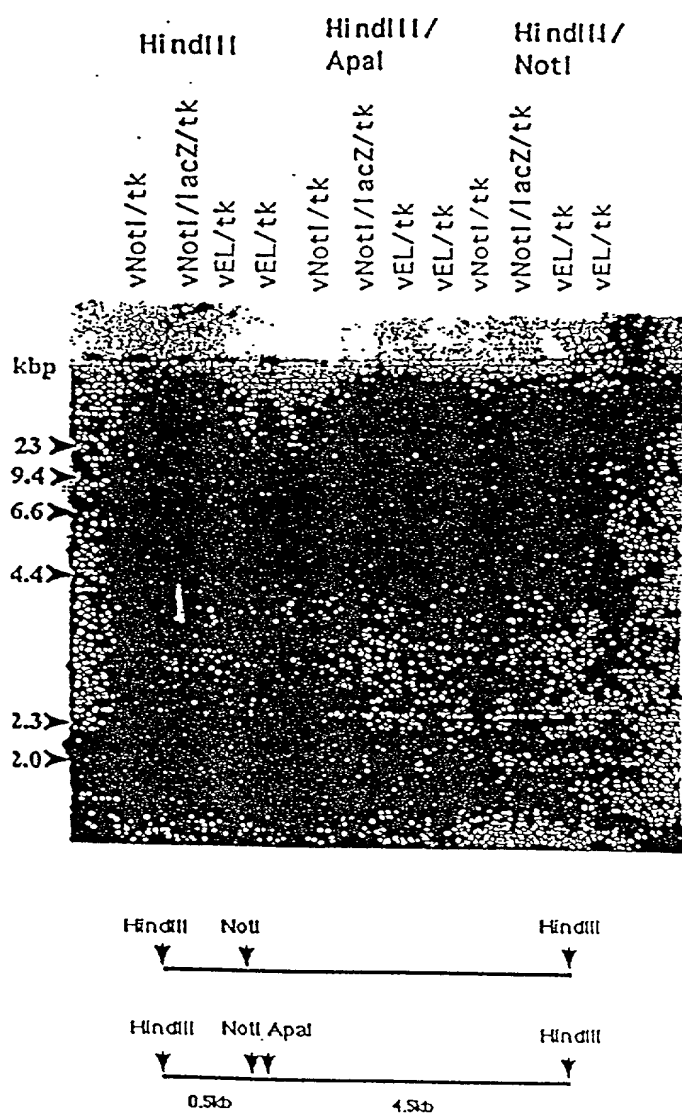


FIGURE 2B

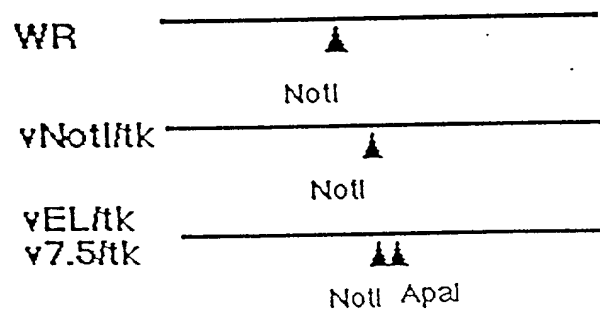
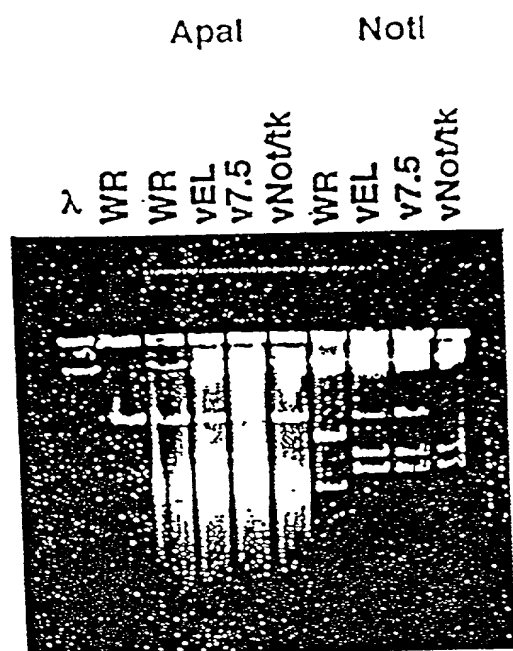


FIGURE 3

Phi X
WR
pJNot
vNotI/tk
vpNotI
p7.5/tk
v7.5/tk
vNotI/lacZ/tk
pEL/tk
vEL/tk

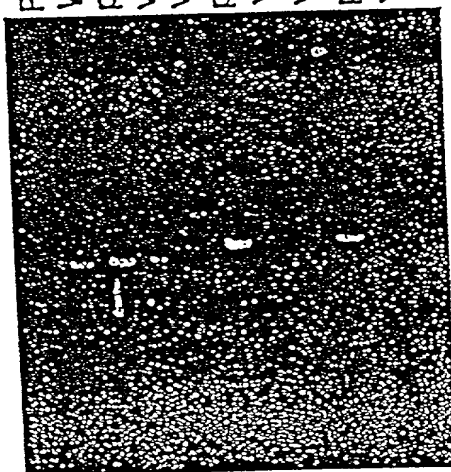


FIGURE 4

B-Glucoronidase Analysis of Recombinant Viruses

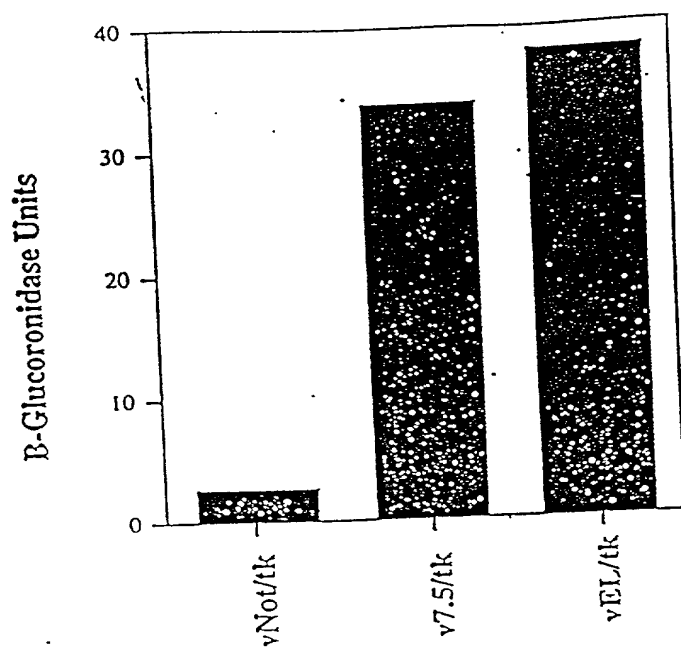


FIGURE 5

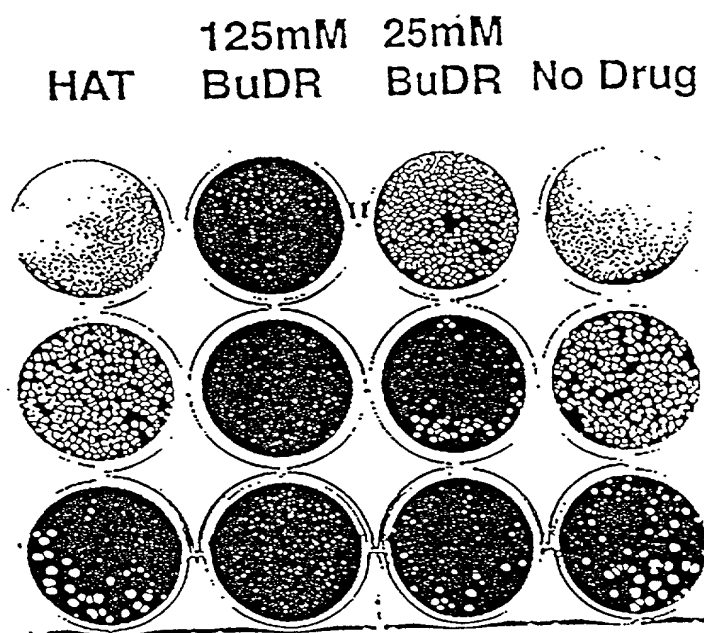


FIGURE 6

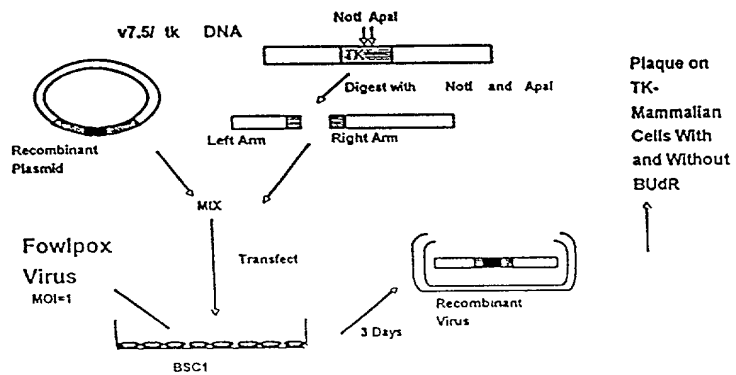


FIGURE 7

1. p7.5tk

7.5K PROMOTER NOTI APAI
5'-GGCCAAAAATTGAAAAACTAGATCTATTATTGCACGGCGGCCGCCATGGGCCCGGCC-3'

2. p7.5/ATG0/tk

7.5K PROMOTER NOTI BAMHI SMAI PSTI
5'-GGCCAAAAATTGAAAAACTAGATCTATTATTGCACGGCGGCCCGGTGATCCCCGGGCTGCAGGAA

TRANSCRIPTION
TRANSLATION STOP CODONS STOP SIGNAL
SALI
TTCGATATCAAGCTTATCGATACCGTCGACCTCGAGGGGGGCCCTAACTAAATTTGTTTGT

APAI
GGGCCCGGCC-3'

3. p7.5/ATG1/th

7.5K PROMOTER	NOTI	START	CODON BAMHI	SMAI	PSTI
5'-GCCCAAATTGAAAACACTAGATCTATTATTGCACGGCGCCGCCATGGTGGATCCCCCGGGCTGCAGGAA					
TTCGATATCAAGCTTATCGATACCGTCGACCTCGAGGGGGGCCCTAACTAATTGTTTTGT					

APAI

'GGGCCCCGGCC-3'

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100

4. p7.5/ATG2/tk

7.5K PROMOTER NOTI START
CODON BAMHI SMAI PSTI

5'-GGCCAAAATTGAAAACTAGATCTATTATTGACGGCGGCCCA TGAGTGGATCCCCCGGCTGCCAGGA

TRANSLATION TRANSCRIPTION
STOP CODONS STOP SIGNAL

SALI

TTCGATATCAAGCTTATCGATACCGTCGACCTCGAGGGGGCCCTAACTAATTTGTTTGT

APAI

GGGCCCCGGCC-3'

5. p7.5/ATG3/tk	START				
7.5K PROMOTER	NOTI	CODON	BAMHI	SMAI	PSTI
5'- GGCCAAAATTGAAAACTAGATCTATTATTGCACGCGCGCCGCCATGACGTGGATCCCCCGGCTGCAGGAA					
		TRANSLATION	TRANSCRIPTION		
		STOP CODONS	STOP SIGNAL		
	SALI				
TTCGATATCAAGCTTATCGATACCGTCGACCTCGAGGGGGGCCTAACTAAATTTGTTTGT					
APAI					
GGGCCCCGGCC-3'					

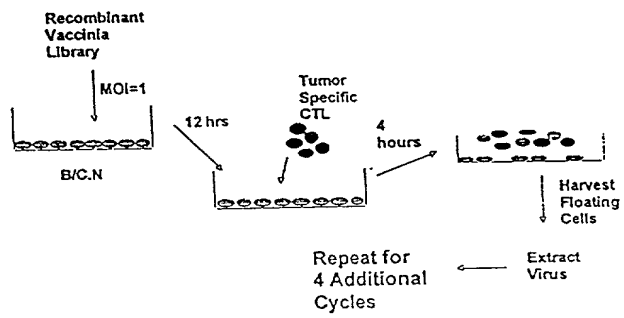


FIGURE 9

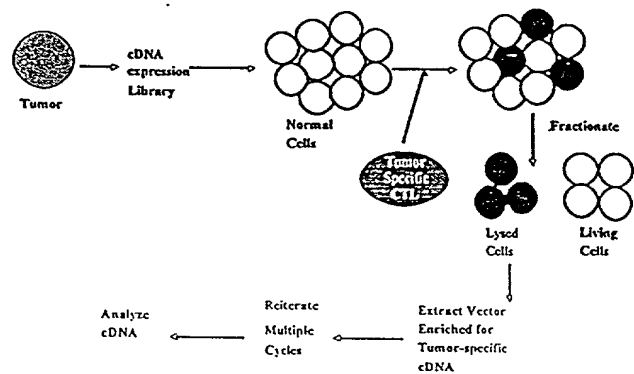


FIGURE 10

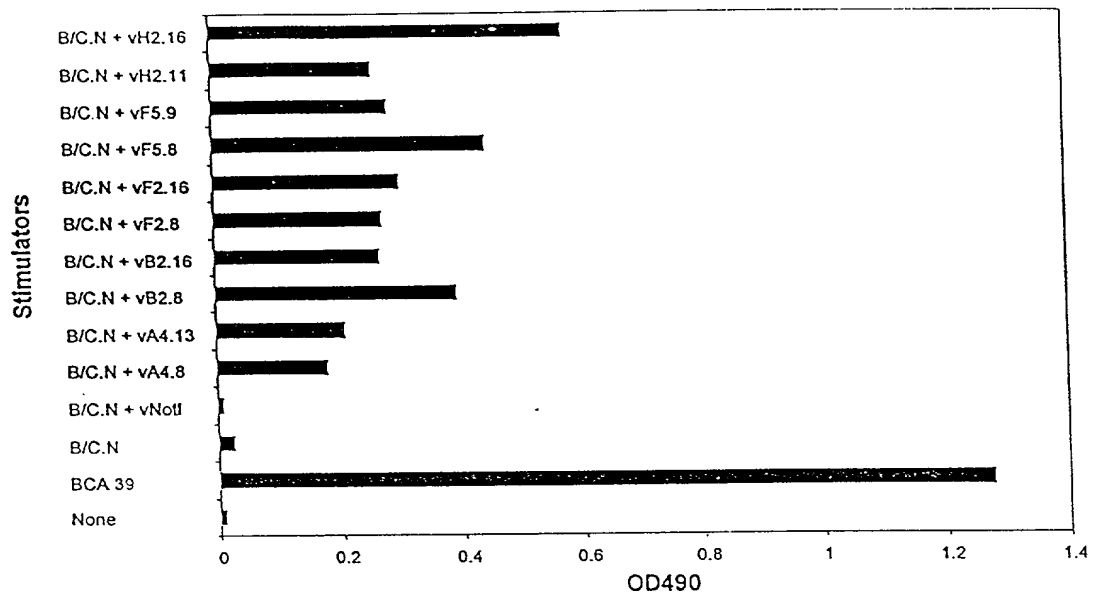


FIGURE IIA

<u>Target</u>	Percent Specific Lysis	
	Effector : Target	
	<u>10:1</u>	<u>2:1</u>
BCA 34	68.4	54.8
BCA 39	36.6	23.4
B/C.N	0.2	0.3
B/C.N + vF5.8	47.5	34.6
B/C.N + vH2.16	67.8	56.2
B/C.N + vaccinia vector	0	0.2

FIGURE 11B

A. L3

Amino Acid Position	45	46	47	48	49	50	51	52	53	54	55	56
Sequence	A	F	L	G	Y	K	A	G	M	T	H	I
Nucleotide	GCC	TTT	CTG	GGT	TAC	AAG	GCT	GGC	ATG	ACC	CAC	ATC

B. H2.16

Amino Acid Position	45	46	47	48	49	50	51	52	53	54	55	56
Sequence	A	F	L	G	Y	K	A	G	M	I	H	I
Nucleotide	---	---	---	---	---	---	---	---	---	-T-	---	---

FIGURE 12

<u>Target</u>	Percent Specific Lysis	
	Effector: Target	
	<u>10:1</u>	<u>2:1</u>
BCA 34	62.4	32.1
BCA 39	49.7	23.6
B/C.N	3.3	0.2
B/C.N + L3 peptide 48-56(I54)	46.0	16.1
B/C.N + L3 peptide 48-56(T54)	2.0	0
B/C.N + L3 peptide 45-54(I54)	0	0

FIGURE 13A

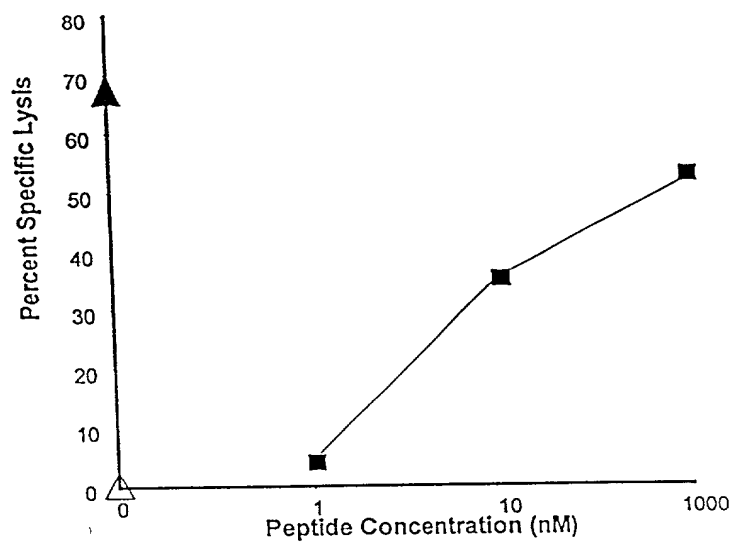
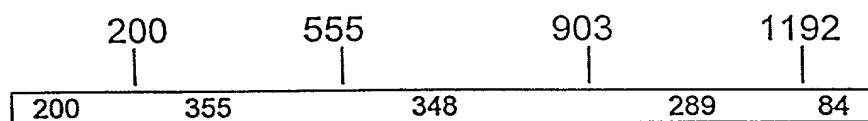


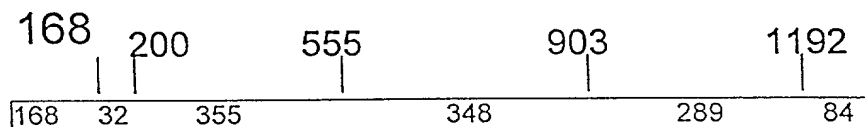
FIGURE 13B

Published L3 (1276 bp)



168-171 = GACC

H2.16 (1276 bp)



168-171 = GATC

FIGURE 14A

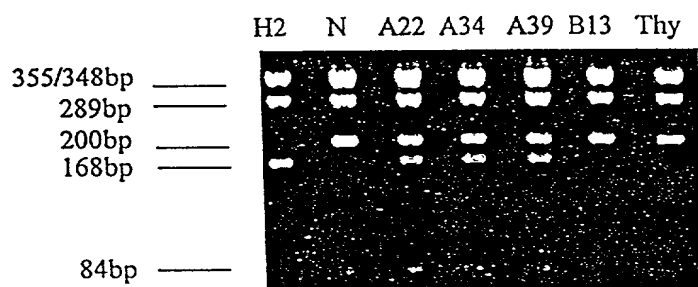


FIGURE 14B

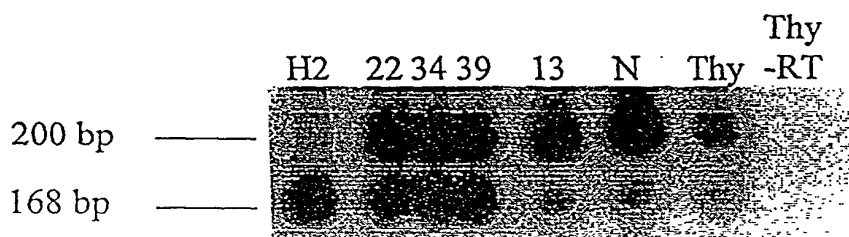


FIGURE 14C

Target	Percent Specific Lysis Immunogen			
	vH2.16		v7.5/tk	
	<u>40:1</u>	<u>10:1</u>	<u>40:1</u>	<u>10:1</u>
BCA 34	33.6	12.9	5.7	4.0
BCA 39	22.1	9.0	5.3	3.1
B/C.N + L3 48-56 (I54)	48.2	20.2	3.9	1.5
B/C.N + L3 48-56 (T54)	6.4	1.4	1.8	2.9
B/C.N	7.1	5.7	6.1	2.8
YAC	1.2	2.5	0	1.8

FIGURE 15 A

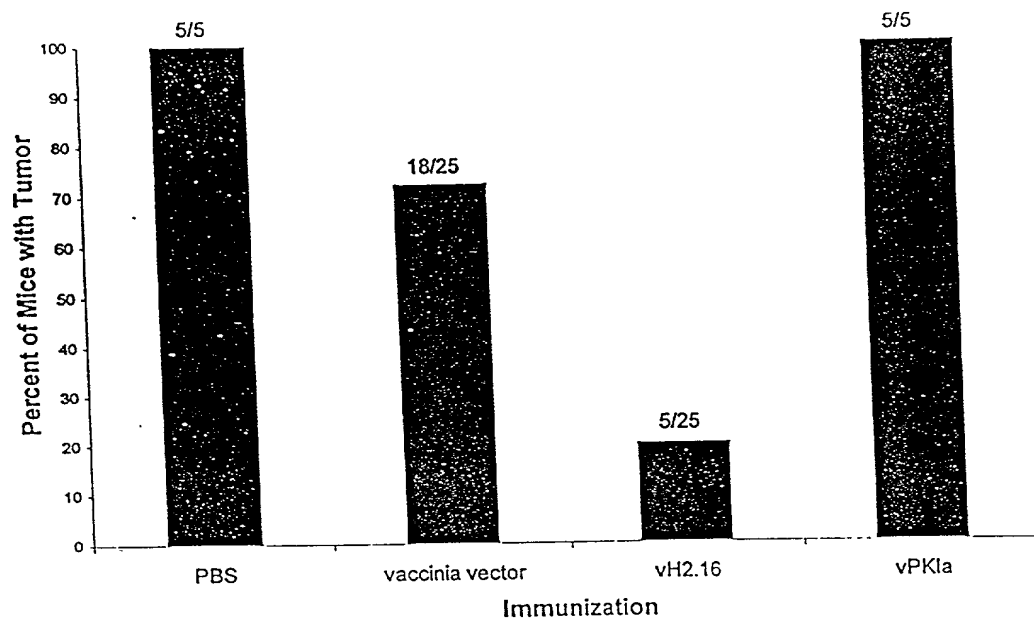


FIGURE 15B

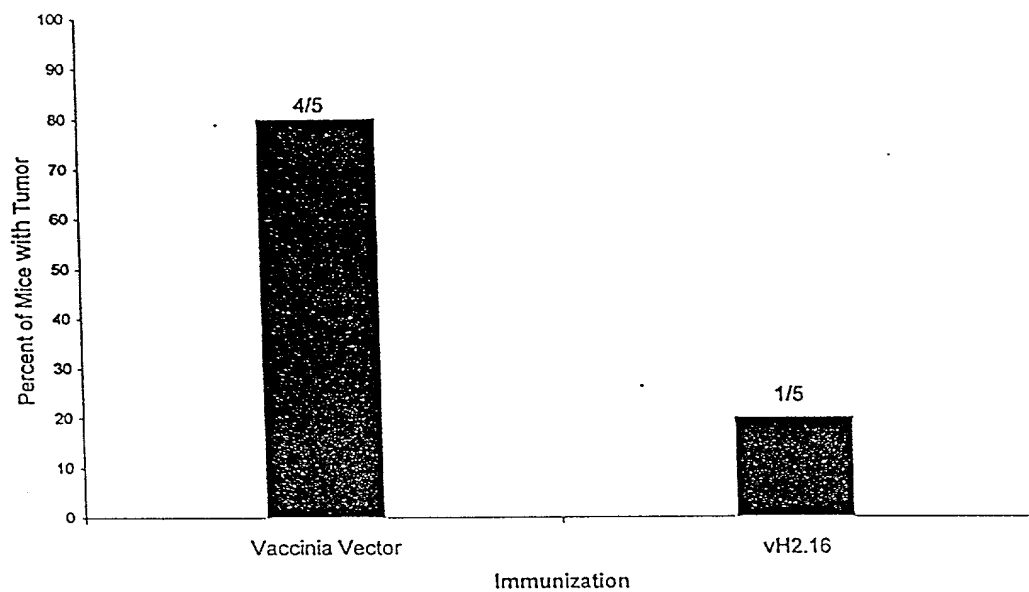


FIGURE 15C

A. Influenza Specific Cytolytic Activity of CD4+
CD45RA+ Human T Cells Stimulated in the Presence
of IL-12 and IL-18.

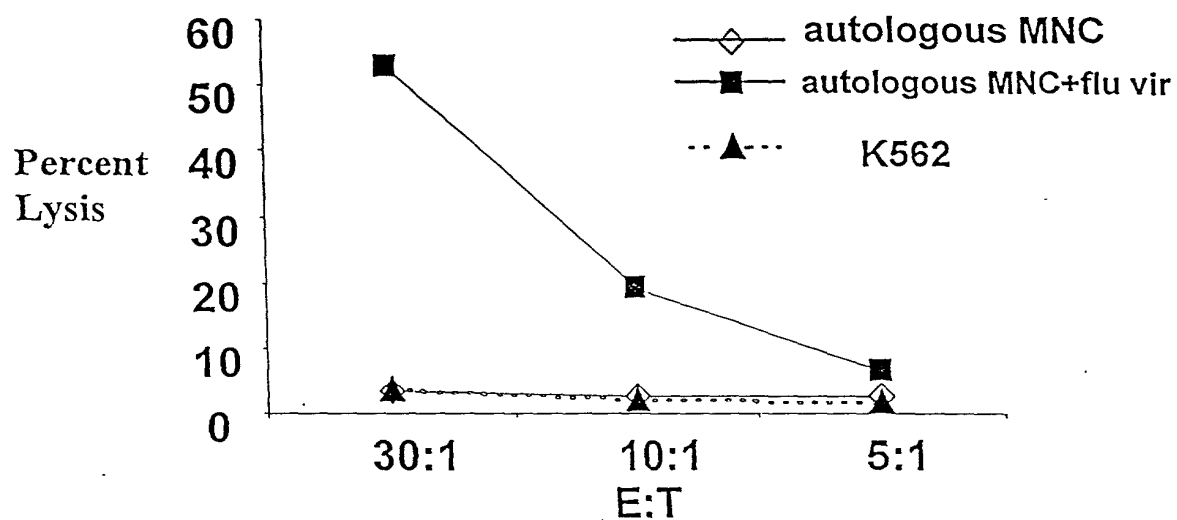


FIGURE 16

CD4+ Primary Cytotoxic T Cell response

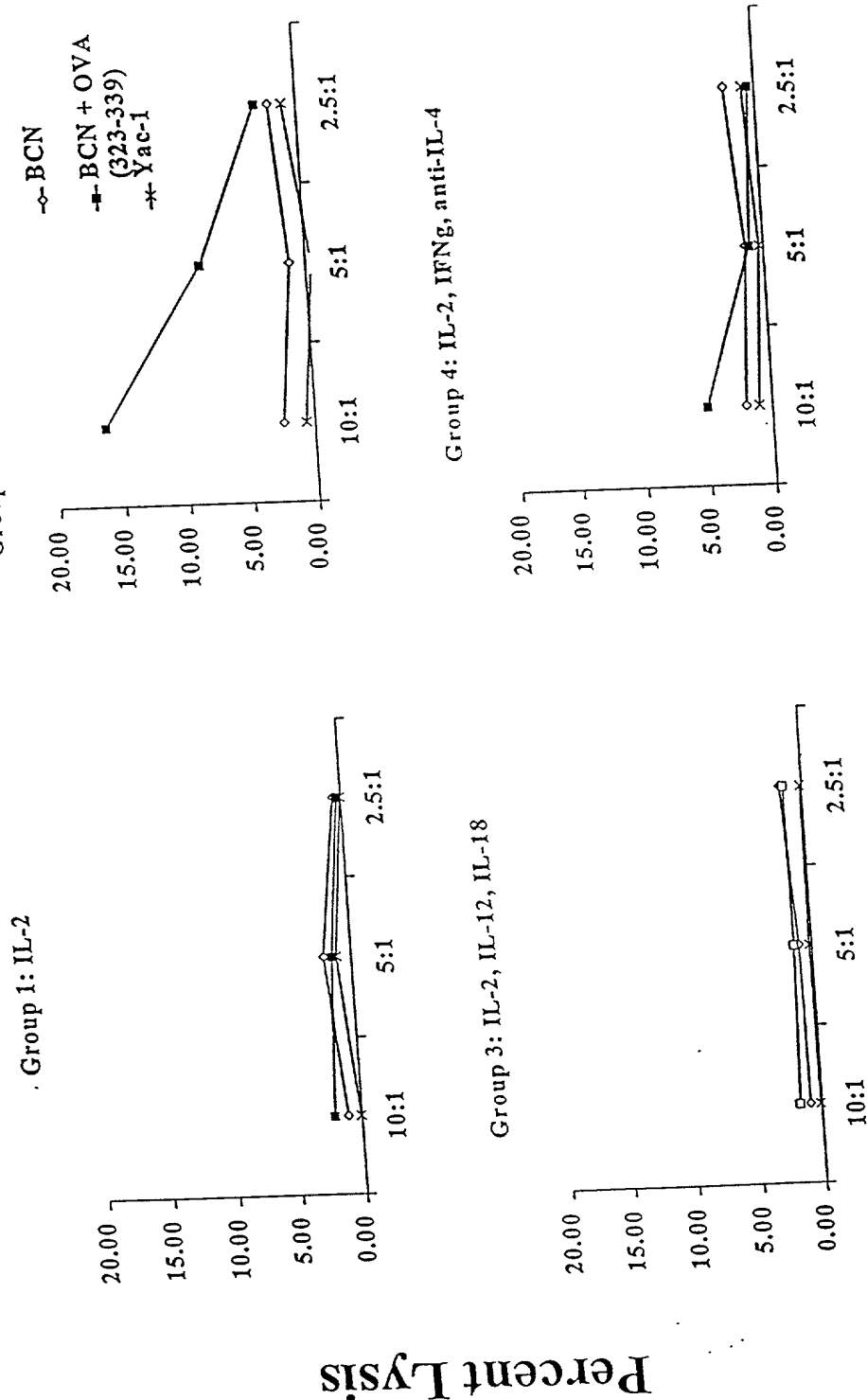


FIGURE 17

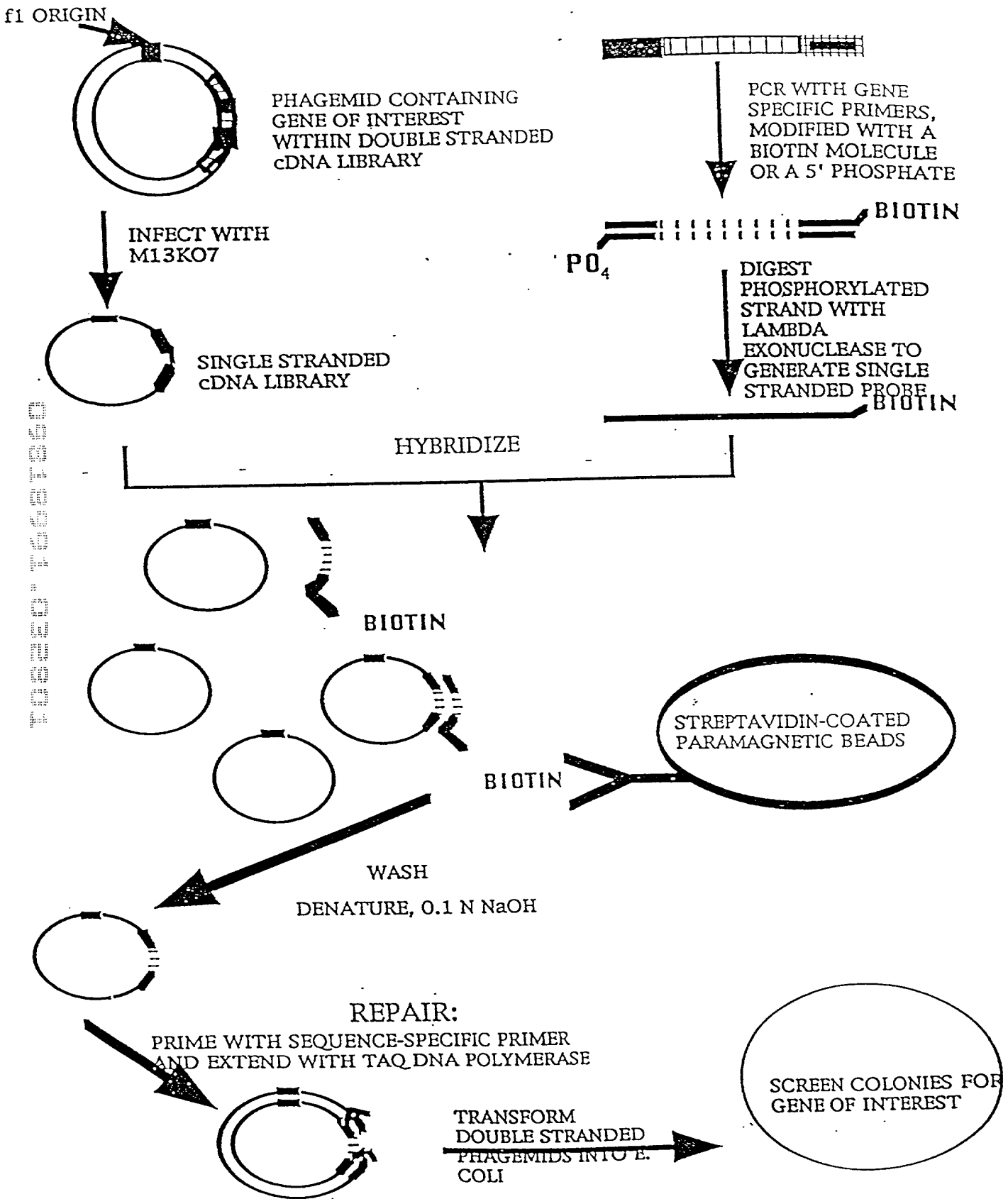


FIGURE 18

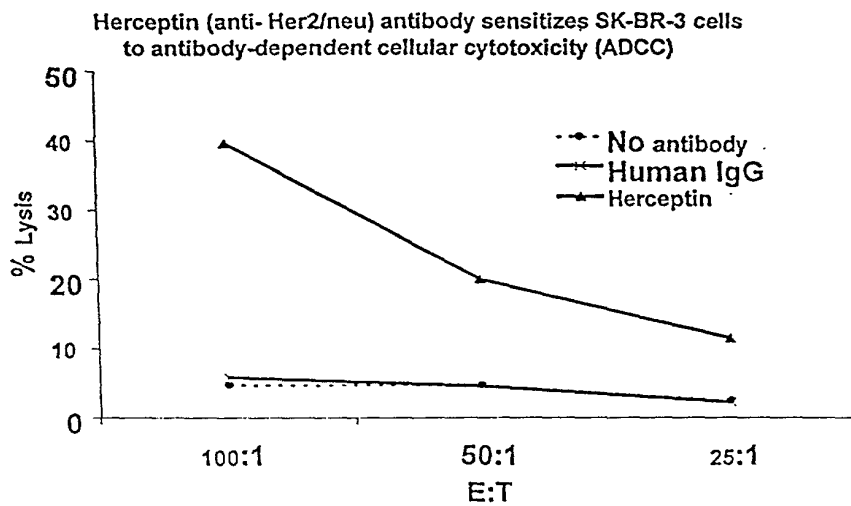


FIGURE 19

Tolerance to Alloantigens Induced in presence of Antigens and Anti-CD40 Ligand Antibody

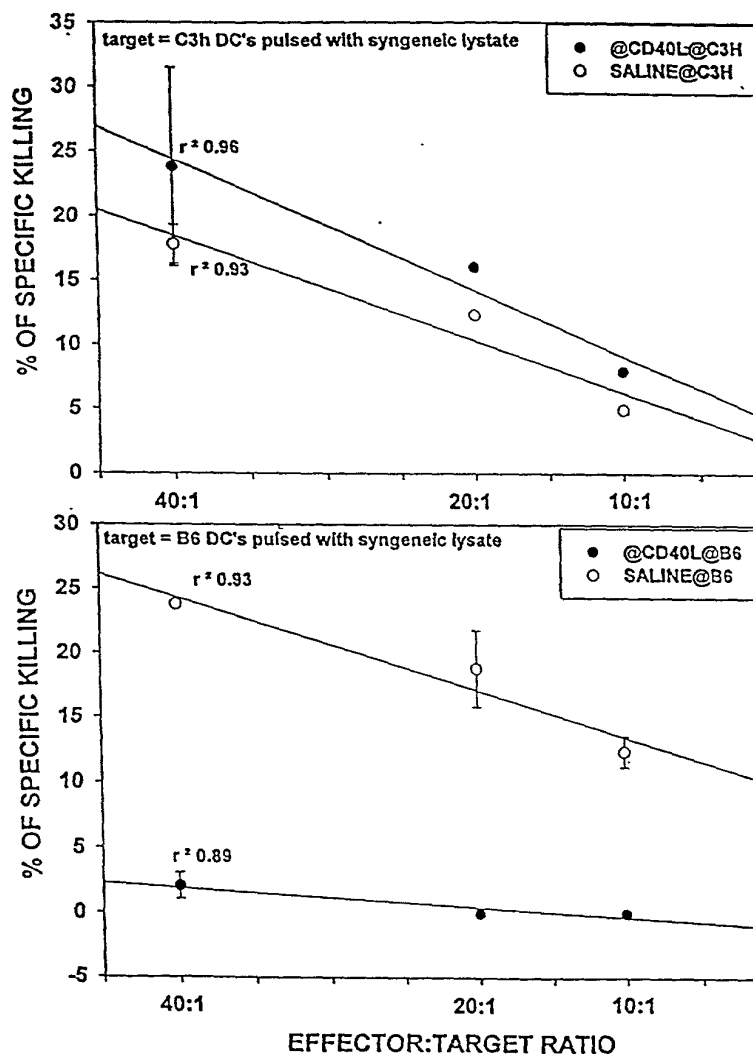


FIGURE 20

Figure 21

